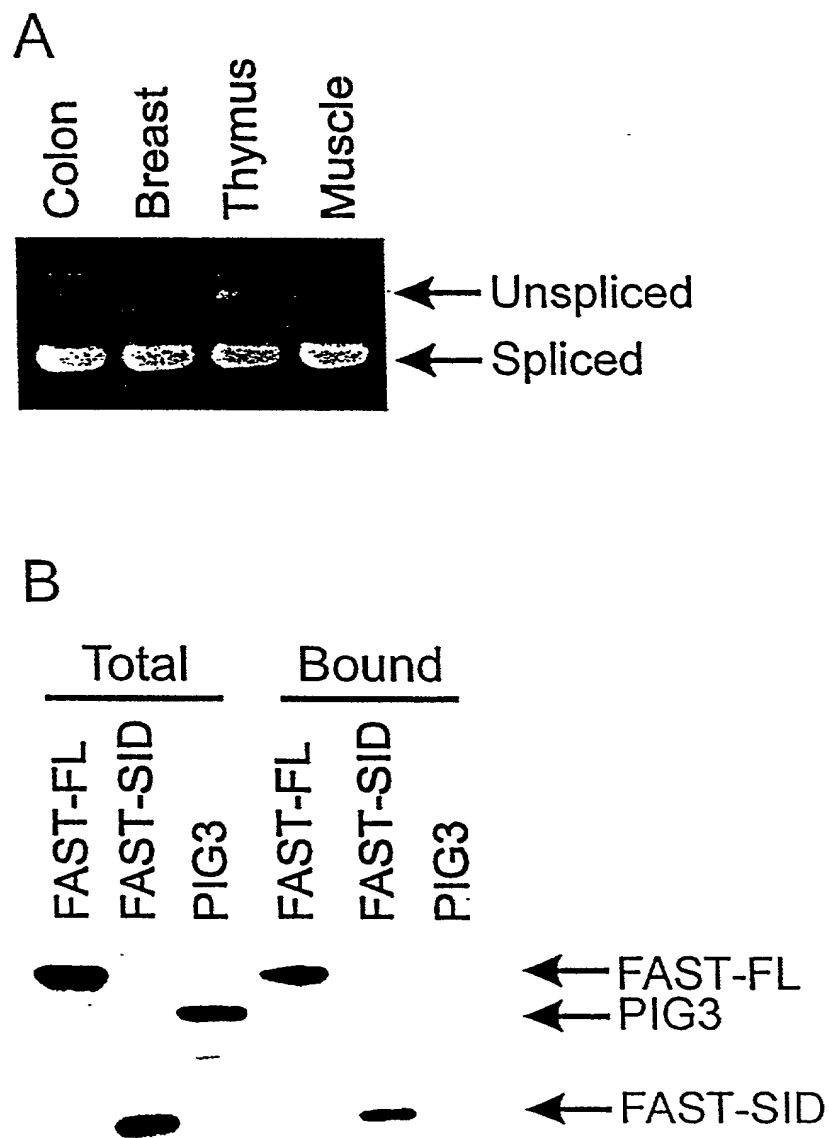


FIG. 1

ST-1	mgpcsgsrlgppaeaspsqpp-----	21
ST-1	mrdpsslysgfagsgyesveppslallssidqeqlpvatggsynhsvqpw	51
ST-1	-----	21
ST-1	pqpwpplslyreggtwspdragsmyglspgthegscththegpkdsmagdht	102
ST-1	---RRRRLRLRHDPPYTYLAMEECAAESRRIRAOIROQAVFPE	69
ST-1	rsrASKKINVHRYYPYSYLAMEEYONSPEKRRLSOLKESTLFPE	153
ST-1	EREDYEGWEDSRHNEENRCHRPKDPAPQAKGNEWANDSLFAEAL	120
ST-1	ENGDMGWEDSRHNESSDCRKLADFGAPQAKGNEWTDYSRELDAM	204
ST-1	REONHACRrwqngCARGAEAKDEGEYELHGRPYrppspppppsegfiks	171
ST-1	KRONHACRTR-----GGSDYEVODLAPYELHNYKvehnagayghqmpgshar	250
ST-1	lggsgegapwpglapqsspvpagtgnsgaeaptpolpsserplwplcpl	222
ST-1	slslaedsqqntntggklntsfmidslldlqendlpdasrnlengrispav	301
ST-1	pgptrvegetvqggaigostlspeprawplhllqgtavpggrssgghrasl	273
ST-1	amnmwssaplllythskotrnarsoglstihstyssssssistispvgfqk	352
ST-1	wgq-----	276
ST-1	eqeksgqrtqrvghpikrsredddcsttssdpdtgnyspieppkmpplsl	403
ST-1	-LPISYLPIYTPNEMElappotscpcpstspaywgvapetrppgllc-	325
ST-1	dlPISTYTKSVAPEMELAPpsvlqffhfrftyynygpspymtppywgfpht	454
ST-1	-----DEDALEFGVPPNKSLDWWVSHPRDEAAFGpggwllsw	362
ST-1	nsggdsprgpgsplDEDNMLRAMPPNKSLDVLTSHEGDEVHESflsqclg	505
ST-1	csl-----	365
ST-1	ssgspypsrqglm	518

FIG. 2



A

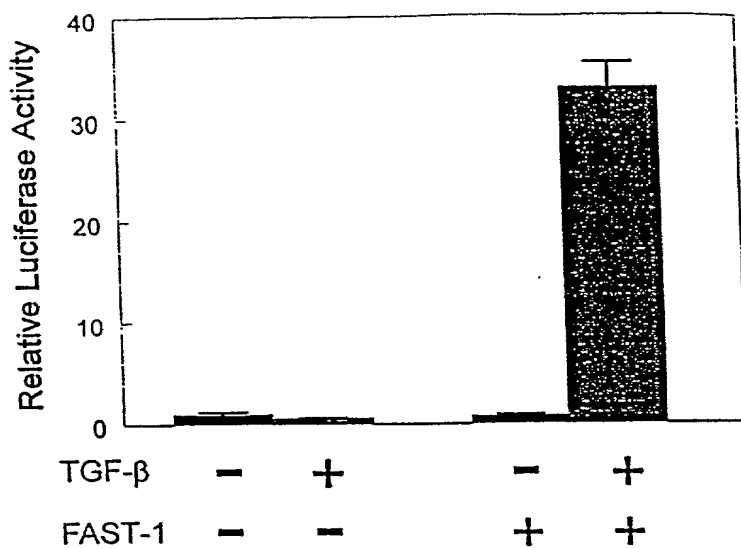
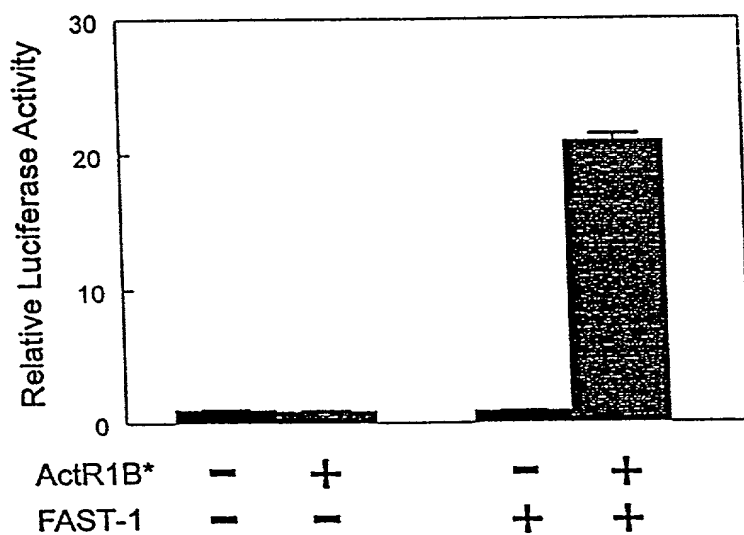


FIG. 3

B



C

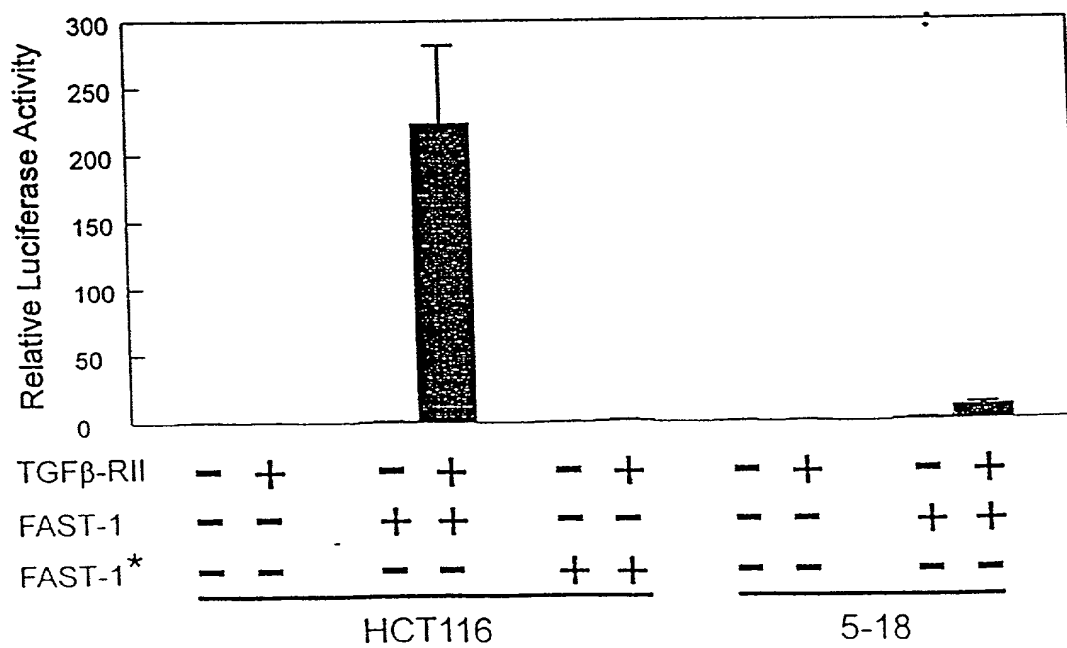


FIG. 4



... CONSENSUS									
	T	G	T	GT	GT	A	T	T	
G	41	41	11	0	100	0	0	0	70 64 52
A	23	35	17	0	0	100	0	0	11 11 17
T	23	23	64	100	0	0	100	100	5 17 0
C	11	0	5	0	0	0	0	0	11 5 29

